LISTING OF CLAIMS

1-23. (Cancelled)

24. (Currently Amended) A method of detecting the presence of antibodies to virulent Myeobacterium M bovis and M. tuberculosis in a biological sample, said method comprising: combining said sample with a protein having the amino acid sequence of SEQ ID NO:2, a Mycobacterial homolog thereof or an antigenic determinant thereof; and detecting antibodies bound to said protein.; wherein said Mycobacterium is M. bovis, M. tuberculosis, M. leprae, M. africanum, M. mieroti, M. avium, M. intracellulare of M. serofulaceum.

- 25. (Cancelled)
- 26. (Previously Added) The method of Claim 24, wherein said protein is immobilized on a solid support.
- 27. (Previously Added) The method of Claim 26, wherein said solid support is nitrocellulose.
- 28. (Previously Added) The method of Claim 24, wherein said sample comprises one or more of sputum, blood, and serum.
- 29. (Previously Added) The method of Claim 24, wherein said detecting is by a qualitative detection system.
- 30. (Previously Added) The method of Claim 29, wherein said qualitative detection system is a horseradish peroxidase-protein A detection system.

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- 31. (Previously Added) The method of Claim 24, wherein said detecting is by a quantitative detection system.
- 32. (Previously Added) The method of Claim 31, wherein said quantitative detection system is a radioimmunoassay.
- 33. (Previously Added) The method of Claim 24, further comprising: combining a control biological sample with said protein; and comparing the detection of said binding to the binding of antibodies in the control sample with said protein.

34-40. (Cancelled)

41. (Previously Amended) A method of detecting the presence of Mycobacterium in a biological sample, said method comprising; lysing the cells in said sample; combining said lysate with antibodies to a protein having the amino acid sequence of SEQ ID NO:2 or an antigenic determinant thereof; and detecting said antibodies bound to protein in said lysate; wherein said Mycobacterium is M. bovis, M. tuberculosis, M. leprae, M. africanum, M. microti, M. avium, M. intracellulare or M. scrofulaceum.

- 42. (Previously Amended) The method of Claim 41, wherein said Mycobacterium is M. bovis.
- 43. (Previously Added) The method of Claim 41, wherein said lysate is immobilized on a solid support.
- 44. (Previously Added) The method of Claim 43, wherein said solid support is nitrocellulose.

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- 45. (Previously Added) The method of Claim 41, wherein said detecting is by a qualitative detection system.
- 46. (Previously Added) The method of Claim 45, wherein said qualitative detection system is a horseradish peroxidase-protein A detection system.
- 47. (Previously Added) The method of Claim 41, wherein said detecting is by a quantitative detection system.
- 48. (Previously Added) The method of Claim 47, wherein said quantitative detection system is a radioimmunoassay.
- 49. (Previously Added) The method of Claim 41, further comprising: culturing a diagnostic sample to produce colonies of bacteria present therein, whereby said culture represents said biological sample.
- 50. (Previously Added) A method of detecting the presence of antibodies to a virulent Mycobacteriam in a biological sample, said method comprising: combining said sample with a purified protein of a mycobacterium other than M. bovis BCG, wherein said protein is a homolog of the protein of SEQ ID NO: 2; is an immunogenic membrane-associated protein of said mycobacterium; and is encoded by DNA which is capable of hybridizing with a DNA probe having the complete sequence represented in SEQ ID NO: 1 under conditions where, on a Southern blot, said probe will identify single 3.25 kb BarnHI fragments from M. bovis BCG and M. tuberculosis H37Rv DNA, but will not hybridize with BarnHI-digested DNA from either M. smegmatis or M. vaccae.
- 51. (Cancelled)

S2.

(Previously Added) The method of Claim 41, wherein said Mycobacterium is M.

tuberculosis.

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